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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/037,311A

DATE: 12/10/2002 P.6

TIME: 15:43:38

Input Set : A:\MS00-001C2 XFTASE.txt

Output Set: N:\CRF4\12102002\J037311A.raw

3 <110> APPLICANT: MICHIGAN STATE UNIVERSITY
 5 <120> TITLE OF INVENTION: XYLOGLUCAN FUCOSYLTRANSFERASES
 7 <130> FILE REFERENCE: MS00-001C2
 9 <140> CURRENT APPLICATION NUMBER: US 10/037,311A
 10 <141> CURRENT FILING DATE: 2001-11-09
 12 <150> PRIOR APPLICATION NUMBER: US60/117,555
 13 <151> PRIOR FILING DATE: 1999-01-28
 15 <160> NUMBER OF SEQ ID NOS: 15
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 558
 21 <212> TYPE: PRT
 22 <213> ORGANISM: Arabidopsis thaliana
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 30 Thr Gly Gly Ser Lys Ser Val Asn Phe Ser Glu Leu Leu Gln Met Lys
 31 20 25 30
 34 Tyr Leu Ser Ser Gly Thr Met Lys Leu Thr Arg Thr Phe Thr Thr Cys
 35 35 40 45
 38 Leu Ile Val Phe Ser Val Leu Val Ala Phe Ser Met Ile Phe His Gln
 39 50 55 60
 42 His Pro Ser Asp Ser Asn Arg Ile Met Gly Phe Ala Glu Ala Arg Val
 43 65 70 75 80
 46 Leu Asp Ala Gly Val Phe Pro Asn Val Thr Asn Ile Asn Ser Asp Lys
 47 85 90 95
 50 Leu Leu Gly Gly Leu Leu Ala Ser Gly Phe Asp Glu Asp Ser Cys Leu
 51 100 105 110
 54 Ser Arg Tyr Gln Ser Val His Tyr Arg Lys Pro Ser Pro Tyr Lys Pro
 55 115 120 125
 58 Ser Ser Tyr Leu Ile Ser Lys Leu Arg Asn Tyr Glu Lys Leu His Lys
 59 130 135 140
 62 Arg Cys Gly Pro Gly Thr Glu Ser Tyr Lys Lys Ala Leu Lys Gln Leu
 63 145 150 155 160
 66 Asp Gln Glu His Ile Asp Gly Asp Gly Glu Cys Lys Tyr Val Val Trp
 67 165 170 175
 70 Ile Ser Phe Ser Gly Leu Gly Asn Arg Ile Leu Ser Leu Ala Ser Val
 71 180 185 190
 74 Phe Leu Tyr Ala Leu Leu Thr Asp Arg Val Leu Leu Val Asp Arg Gly
 75 195 200 205
 78 Lys Asp Met Asp Asp Leu Phe Cys Glu Pro Phe Leu Gly Met Ser Trp
 79 210 215 220
 82 Leu Leu Pro Leu Asp Phe Pro Met Thr Asp Gln Phe Asp Gly Leu Asn

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87					245					250						255
90	Asp	Thr	Glu	Gly	Thr	Leu	Ser	His	Leu	Tyr	Leu	His	Leu	Val	His	Asp
91						260				265						270
94	Tyr	Gly	Asp	His	Asp	Lys	Met	Phe	Phe	Cys	Glu	Gly	Asp	Gln	Thr	Phe
95						275				280						285
98	Ile	Gly	Lys	Val	Pro	Trp	Leu	Ile	Val	Lys	Thr	Asp	Asn	Tyr	Phe	Val
99						290				295						300
102	Pro	Ser	Leu	Trp	Leu	Ile	Pro	Gly	Phe	Asp	Asp	Glu	Leu	Asn	Lys	Leu
103						305				310			315			320
106	Phe	Pro	Gln	Lys	Ala	Thr	Val	Phe	His	His	Leu	Gly	Arg	Tyr	Leu	Phe
107							325				330					335
110	His	Pro	Thr	Asn	Gln	Val	Trp	Gly	Leu	Val	Thr	Arg	Tyr	Tyr	Glu	Ala
111						340				345						350
114	Tyr	Leu	Ser	His	Ala	Asp	Glu	Lys	Ile	Gly	Ile	Gln	Val	Arg	Val	Phe
115						355				360						365
118	Asp	Glu	Asp	Pro	Gly	Pro	Phe	Gln	His	Val	Met	Asp	Gln	Ile	Ser	Ser
119						370				375						380
122	Cys	Thr	Gln	Lys	Glu	Lys	Leu	Leu	Pro	Glu	Val	Asp	Thr	Leu	Val	Glu
123						385				390			395			400
126	Arg	Ser	Arg	His	Val	Asn	Thr	Pro	Lys	His	Lys	Ala	Val	Leu	Val	Thr
127							405				410					415
130	Ser	Leu	Asn	Ala	Gly	Tyr	Ala	Glu	Asn	Leu	Lys	Ser	Met	Tyr	Trp	Glu
131						420				425						430
134	Tyr	Pro	Thr	Ser	Thr	Gly	Glu	Ile	Ile	Gly	Val	His	Gln	Pro	Ser	Gln
135						435				440						445
138	Glu	Gly	Tyr	Gln	Gln	Thr	Glu	Lys	Lys	Met	His	Asn	Gly	Lys	Ala	Leu
139						450				455						460
142	Ala	Glu	Met	Tyr	Leu	Leu	Ser	Leu	Thr	Asp	Asn	Leu	Val	Thr	Ser	Ala
143						465				470			475			480
146	Trp	Ser	Thr	Phe	Gly	Tyr	Val	Ala	Gln	Gly	Leu	Gly	Gly	Leu	Lys	Pro
147							485				490					495
150	Trp	Ile	Leu	Tyr	Arg	Pro	Glu	Asn	Arg	Thr	Thr	Pro	Asp	Pro	Ser	Cys
151							500				505					510
154	Gly	Arg	Ala	Met	Ser	Met	Glu	Pro	Cys	Phe	His	Ser	Pro	Pro	Phe	Tyr
155						515				520						525
158	Asp	Cys	Lys	Ala	Lys	Thr	Gly	Ile	Asp	Thr	Gly	Thr	Leu	Val	Pro	His
159						530				535						540
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174	aagtccgtta	atttctccga	actacttcaa	atgaagtatc	ttagctccgg	tacgatgaag										120
176	ctcacgagaa	ctttcactac	ttgcttgata	gtcttctctg	tactagtagc	attctcaatg										180

RAW SEQUENCE LISTING
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178	atctttcacc	aacacccatc	tgattcaaat	cggatttatgg	gtttcgccga	agcttagagtt	240
180	ctcgacgccc	gagtttccc	aaattctgat	aagcttctcg	gagggtact	tgcttctgggt	300
182	tttgatgaag	attcttgcct	tagtaggtac	caatcagttc	attaccgtaa	actttcaccc	360
184	tacaaggccat	cttcttatct	catctctaag	cttagaaaact	acgaaaagct	tcacaagcga	420
186	tgtggtccgg	gtactgaatc	ttacaagaaa	gctctaaaac	aacttgtatca	agaacatatt	480
188	gatggtgatg	gtgaatgcaa	atatgttgg	tggatttctt	ttagcggctt	aggaacagg	540
190	atactttctc	tagcctcggt	tttctttac	gcmccttaa	cgatagagt	cttgcttgg	600
192	gaccgagggaa	aagacatgga	tgatctctt	tgcgagccgt	ttctcggtat	gtcggttgt	660
194	ctacctttag	attcccttat	gactgatcag	tttgatggat	taaatcaaga	atcatctcg	720
196	tgttatggat	atatggtcaa	gaatcaggtt	attgatactg	agggaaacctt	gtctcatctt	780
198	tatcttcatc	ttgttcatga	ttatggagat	catgataaga	tgttcttctg	tgaaggagac	840
200	caaacattca	tcgggaaaagt	cccttgggtt	attgttaaaa	cagacaatta	cttgttcca	900
202	tctctgtgg	taataccggg	tttcgatgat	gaactaaaca	agctattccc	acagaaagcgt	960
204	actgtcttc	atcaacttgg	taggtatctt	tttcacccaa	ctaaccagg	atggggctt	1020
206	gtcactagat	actacgaagc	ttacttatacg	catcggtat	agaagatgg	gattcaagta	1080
208	agagtttcg	atgaagaccc	gggtccattt	cagcatgtga	tggatcagat	ttcatcttgc	1140
210	actcaaaaag	agaaacttct	acctgaagta	gacacactag	tggagagatc	tcgccccatgtt	1200
212	aataccccca	aacacaaaagc	cgtgcttgc	acatcttga	acgcgggtt	cgcggagaac	1260
214	ttaaagagta	tgtattggg	atatccgaca	tcaactggag	aaatcatcg	tgttcatcag	1320
216	ccgagccaaag	aagggttatca	gcagaccgaa	aaaaagatgc	ataatggca	agctcttgc	1380
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220	tatgttagctc	aaggcttgg	aggtttaaag	ccttggatac	tctatagacc	cgaaaaccgt	1500
222	acaactcccg	atccttcgt	ttgtcgggct	atgtcgatgg	agccttgc	ccacttcgc	1560
224	ccattctatg	attgtaaaagc	gaaaacgggt	attgacacgg	gaacactagt	tcctcatgt	1620
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241	atcttcacc	aacacccatc	tgattcaaat	cggatttatgg	gtttcgccga	agcttagagtt	240
243	ctcgacgccc	gagtttccc	aatgttact	aacatcagta	tgtgttctt	caagtcaaag	300
245	tttgagctt	tattacttta	gatctcgatc	tttacactac	gcatttgcct	ctgtatgtc	360
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253	tttggtttt	tagattctga	taagcttctc	ggagggtac	ttgcttctgg	ttttgatgaa	600
255	gattcttgc	ttagtaggt	ccaatcagg	cattaccgt	aaccttca	ttacaagcca	660
257	tcttcttata	tcatctctaa	gcttagaaac	tacgaaaagc	ttcacaagcg	atgtggtcc	720
259	ggtactgaat	cttacaagaa	agctctaaa	caacttgc	aagaacat	tgtatgtat	780
261	ggtgaatgca	aatatgtt	gtggatttct	tttagcggct	tagggaaacag	gatactttct	840
263	ctagcctcg	ttttcttta	cgcgtttt	acggatagag	tcttgcgt	tgaccgaggg	900
265	aaagacatgg	atgatcttt	ttgcgagcc	tttctcggt	tgtcggtgtt	gctacctt	960
267	gattcccta	tgactgatca	tttgatgga	ttaaatcaag	aatcatctcg	ttgttatg	1020
269	tatatggta	agaatcagg	gattgatact	gagggaaactt	tgtctcatct	ttatcttcat	1080
271	cttgcgtat	attatqqaga	tcatgataaq	atgttcttct	gtqaaggqaa	ccaaacattc	1140

RAW SEQUENCE LISTING

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277	catcaacttag gtaggtatct tttcaccca actaaccaag tatggggctt agtcactaga	1320
279	tactacgaag cttacttatac gcatgcggat gagaagattg ggattcaagt aagagtttc	1380
281	gatgaagacc cgggtccatt tcagcatgtg atggatcaga tttcatctt tactcaaaaa	1440
283	gagaaacttc tacctgaagt agacacacta gtggagagat ctcgccatgt taatacccc	1500
285	aaacacaaag ccgtgcttgt cacatcttg aacgcgggtt acgcggagaa cttaaagagt	1560
287	atgtattggg aatatccgac atcaacttggaa gaaatcatcg gtgttcatca gccgagccaa	1620
289	gaaggttatc agcagaccga aaaaaagatg cataatggca aagctcttc gaaaaatgtat	1680
291	ctttttaggt tgacagataa tcttgcata agtgcttggt ctacatttgg atatgttagct	1740
293	caaggtcttggagttaaa gccttggata ctctatagac cccaaaaccg tacaactccc	1800
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312	atgaaaccaa atgtgacatc ctcatacgat gattcatcac tagtgcagag agatcaagaa	180
314	caaaaaggta aacttacttt cttcttttg ttttggaaatg tttctaaatt tttctttgaa	240
316	tgtttcatca gatttgcata atatgtctt gcttggaggg ctacttgcgtt ctgggttcaa	300
318	gaaagagtct tgcttgcata gataccatc ttacctctac cgtaaagctt caccgtataa	360
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322	accggaaaca agacagtata ccaatgcaga aagattgctt aaacagaaaac aaacaggta	480
324	gatgaaatca caaggatgca agtatgttg ttggatgtcg ttttagcggat taggaaacag	540
326	gattatcagt attgcttctg ttttgcata tgcaatgttgc acagatagag tcttgcttgc	600
328	tgaaggaggg gaacagttcg cggattttt ctgcgaaccg ttccctgcata ccacctgggtt	660
330	actaccgaaa gatttcacct tagcttagtca gttcagtggc tttggtcaaa actcagctca	720
332	ctgccatgga gatatgtca agaggaaact gattaatgaa tccctctgtt cgtctctgtc	780
334	tcatctctat cttcatcttag ctcatacgacta caatgagcac gacaaaatgt tcttgcata	840
336	agaagatcaa aatctctttaa agaatgttcc ttgggtgatc atgaggacaa acaacttctt	900
338	tgcaccgtct cttttcttgc tttcttctt cgaagaagag ctcggatata tgtttcccga	960
340	gaaaggaacg gtttttcacc atttaggtcg ttacctttc catccttcaa atcaagtctg	1020
342	gggactaatac acaagatact atcaagctt cttagccaaa gctgatgaaa ggattggct	1080
344	tcaaataaga gttttgtatc agaaaatccgg cgtatctctt cgagtcacaa agcaaatcat	1140
346	ttcgtgtgtt caaaaacgaga atctgttacc gagactaagc aaaggtgaag aacaatacaa	1200
348	gcagccatca gaagaagagt tgaaaactcaa atctgtctt gtcacccctt taacaacagg	1260
350	atactttgatc atcttgcataa caatgtattt gggaaaatcca actgtacaa gagatgtat	1320
352	tggaaatcatc cagccaaatgc atgaaggacata ccaacaaaaca gagaagctaa tgcataacag	1380
354	gaaagcttgg gcagagatgtt acttactcgtt cttaaacggat aagttggta ttatgtcttgc	1440
356	gtctacattt gtttatgtatc tcaaggact tggaggatc agagcttggta ttctgtataa	1500
358	acaagagaat caaaccacca accaaatccacc ttgcggtaga gctatgtcac cagatccttgc	1560
360	tttccatgtc cttccatctt atgattgcata agcaaaagaaa ggaactgaca ctggtaatgt	1620
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367	<210> SEQ ID NO: 5	

RAW SEQUENCE LISTING
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 369 <212> TYPE: PRT
 370 <213> ORGANISM: Arabidopsis thaliana
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 379 20 25 30
 382 Val Gln Ala Ser Arg Phe Ile Thr Met Glu Pro Asn Val Thr Ser Ser
 383 35 40 45
 386 Ser Asp Asp Ser Ser Leu Val Gln Arg Asp Gln Glu Gln Lys Asp Ser
 387 50 55 60
 390 Val Asp Met Ser Leu Leu Gly Gly Leu Leu Val Ser Gly Phe Lys Lys
 391 65 70 75 80
 394 Glu Ser Cys Leu Ser Arg Tyr Gln Ser Tyr Leu Tyr Arg Lys Ala Ser
 395 85 90 95
 398 Pro Tyr Lys Pro Ser Leu Leu Ser Lys Leu Arg Ala Tyr Glu Glu
 399 100 105 110
 402 Leu His Lys Arg Cys Gly Pro Gly Thr Arg Gln Tyr Thr Asn Ala Glu
 403 115 120 125
 406 Arg Leu Leu Lys Gln Lys Gln Thr Gly Glu Met Glu Ser Gln Gly Cys
 407 130 135 140
 410 Lys Tyr Val Val Trp Met Ser Phe Ser Gly Leu Gly Asn Arg Ile Ile
 411 145 150 155 160
 414 Ser Ile Ala Ser Val Phe Leu Tyr Ala Met Leu Thr Asp Arg Val Leu
 415 165 170 175
 418 Leu Val Glu Gly Gly Glu Gln Phe Ala Asp Leu Phe Cys Glu Pro Phe
 419 180 185 190
 422 Leu Asp Thr Thr Trp Leu Leu Pro Lys Asp Phe Thr Leu Ala Ser Gln
 423 195 200 205
 426 Phe Ser Gly Phe Gly Gln Asn Ser Ala His Cys His Gly Asp Met Leu
 427 210 215 220
 430 Lys Arg Lys Leu Ile Asn Glu Ser Ser Val Ser Ser Leu Ser His Leu
 431 225 230 235 240
 434 Tyr Leu His Leu Ala His Asp Tyr Asn Glu His Asp Lys Met Phe Phe
 435 245 250 255
 438 Cys Glu Glu Asp Gln Asn Leu Leu Lys Asn Val Pro Trp Leu Ile Met
 439 260 265 270
 442 Arg Thr Asn Asn Phe Phe Ala Pro Ser Leu Phe Leu Ile Ser Ser Phe
 443 275 280 285
 446 Glu Glu Glu Leu Gly Met Met Phe Pro Glu Lys Gly Thr Val Phe His
 447 290 295 300
 450 His Leu Gly Arg Tyr Leu Phe His Pro Ser Asn Gln Val Trp Gly Leu
 451 305 310 315 320
 454 Ile Thr Arg Tyr Tyr Gln Ala Tyr Leu Ala Lys Ala Asp Glu Arg Ile
 455 325 330 335
 458 Gly Leu Gln Ile Arg Val Phe Asp Glu Lys Ser Gly Val Ser Pro Arg
 459 340 345 350
 462 Val Thr Lys Gln Ile Ile Ser Cys Val Gln Asn Glu Asn Leu Leu Pro

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 12/10/2002
PATENT APPLICATION: US/10/037,311A TIME: 15:43:39

Input Set : A:\MS00-001C2 XFTASE.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 10,29,61,92,101,133,147,168,197,215
Seq#:7; Xaa Pos. 10,20,31,34,49,56,66,72
Seq#:8; N Pos. 4,29,146,190,195,224,263,354,382,383,397,408,426,433,434,438
Seq#:8; N Pos. 454,481,489,511
Seq#:11; N Pos. 148,150,221,248,330,382,410,422,451,502,509,528,539,549,647
Seq#:11; N Pos. 650,659,701,702
Seq#:12; N Pos. 276,361,386,409,433,481,490
Seq#:15; N Pos. 4,5,12,16,27,50,53,72,73,81,87,98,118,147,153,159,168,205
Seq#:15; N Pos. 275

VERIFICATION SUMMARY

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L:576 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:60
L:580 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:120
L:582 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:180
L:642 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:16
L:650 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:32
L:654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:48
L:658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:64
L:780 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:784 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:120
L:786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:180
L:788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:240
L:790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:300
L:792 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:360
L:794 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:420
L:796 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:480
L:979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:120
L:981 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:180
L:983 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:240
L:985 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:300
L:987 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:360
L:989 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:420
L:991 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:480
L:993 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:540
L:995 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:600
L:997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:660
L:1060 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:240
L:1064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:360
L:1066 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:420
L:1068 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:480
L:1211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:1213 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:60
L:1215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:120
L:1217 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:180
L:1219 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:240